

Supplementary figure for

Collagen V is potential substrate for clostridial collagenase G in pancreatic islet isolation

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Col-V α 1 sequence

MDVHTRWKARSALRPGAPLLPPLLLLWAPPSRAAQPADLLKVLDHFHNLPDGITKTTGFCATRR
SSKGPDVAYRTKDAQLSAPTKQLYPASAFAFPEDFSILTVKAKKGSQAFLVSIFYNEQQIGLEL
GRSPVFLYEDHTGKPGPEDYPLFRGINLSDGKWHRIALSVHKKNVTLLIDCKKKTFLDRSDHPM
IDINGIIVFGTRILDEEVFEGDIQQLLFVSDHRAAYDYCEHYSPDCDTAVPDTPQSQDPNPDEYYT
EGDGEGETYYEYPYYEDPEDLGKEPTPSKKPVEAAKETTEVPEELTPPTEAAPMPETSEGAGKE
EDVGIGDYDVPSEDYYTPSPYDDLTYGEGEENPDQPTDGGAGAEIPTSTADTSNSSNPAPPGE
ADDLEGEFTEETIRNLDENYYDPYYDPTSSPSEIGPGMPANQDTIYEGIGGPRGEKGQKGEPAIE
PGMLIEGPPGPEGPAGLPGPPGTMGPTQVGDGERGPPGRPGLPGADGLPGPPGTMLMLPFRFGG
GSDAGSKGPMVSAQESQAQAILQQARLALR

GPAGPMGLTGRPGPVGPPGGGLKGEPGDVGPGQGPRGVQGPPGPAGKPGRRGRAGSDARGMPGQT
GPKGDRGFDGLAGLPGEKGHRGDPGSGPPGPGDDGERGDDGEVGPRLPGEPGPRGLLGPKGPP
GPPGPVGVTGMDGQPGPKGNVGPQGEPGPPQQGNPGAQGLPGPQGAIGPPGEKGPLGKPLGPMP
GADGPAGHPGKEGPPGEKGGQGPQGPQGPIGYPGPRGVKGADGIRGLKGTEKGEDGFPGFKDM
GIKGDRGEIGPPGPRGEDGPEGPKGRGGPNGDPGLGPPGEKGKLGVPGPLPGYGRQGPKGSIGFP
GFPGANKEKGGRTPGKPGPRGQRGPTGPRGERGPRGITGKPGPKGNSGDGPAGPPGERGPNGPQ
GPTGFPGPKGPGPPGKDGLPGHPQRGKTFQGKTGPPGPPGVVGPQGPGETGPMERGHPGPP
GPPGEQGLPGLAGKEGTKGDPGPAGLPGKDGPGLRGFPGDRGLPGPGALGLKGNEGPPGPPGA
GSPGERGPAGAAGPIGIPGRGPQGPAGEKGAPGEKGPGPAGRDGLQGPVGLPGPAGPVGPP
GEDGDKGEIGEPQKGSKGDKGEQGPPTGPQGPIGQPGPSGADGEPGPRGQQGLFGQKGDEGPR
GFPGPAGPVGQLQGLPGPPGEKGETGDVGQMGPPGPPGRGPSGAPGADGPQGPPGGIGNPGAVGEK
GEPGEAGEPGLPGEGGPPGPKGERGEKGESGPGSAAGPPGPKPPGDDGPKGSPPVGFPGDPGPP
GEPGPAGQDGPPGDKGDDGEPGQTGSPGPTGEPGPSGPPGKRGPPGAGPEGRQGEKGAKGEAGLE
GPPGKTGPIGPQGAPGKPGPDGLRGIPGPVGEQGLPGSPGPDPGPPGMPPGLPGLKGDSGPKEK
GHPGLIGLIGPPGEQGEKGDRGLPGQGSSGPKGEQGITGPSGPIGPAGPGLPGPPGPKGAKGSS
GPTGPKGEAGHPGPPGPPGEV

IQPLPIQASRTRRNIDASQLDDGNGENYVDYADGMEEIFGSLNSLKEIEQMKRPLGTQQNPART
CKDLQLCHPDFDGEYWVDPNQGCSRDSFKVYCNFTAGGSTCVFPDKKSEGARITSWPKENPGSWF
SEFKRGKLLSYVDAEGNPVGVVQMTFLRLLSASAHQNVTYHCYQSVAWQDAATGSYDKALRFLGSN
DEMSYDNNPYIRALVDGCATKKGYQKTVLEIDTPKVEQVPIVDIMFDFGEASQKFGFEVGPACF
MG

Col-V α2 sequence

MMANWAEARPLLILIVLLGQFVSIKAQEEDDEGYGEEIACTQNGQMYLNNDIWKPAPCQICVCDN
GAILCDKIECDVLDCADPVTGGCCPVCQTPGGNTNFGRGRKGQKGEPLVPVVTGIRGRPG
PAGPPGSQGPRGERGPGRGPQGIDGEPGVPGQPGAPGPPGHPSHPGDLSRPFSAQMAGL
DEKSGLGSQVGLMP

GSVGPVGPRGPQGLQQGGAGPTGPPGEPGDPGPMGPIGSRGPEGPPGKPGEDGEPGRNGNPGEV
GFAGSPGARGFPGAPGLPGLKGHRGHKGLEGPKGEVGAPSKGEAGPTGPMGAMGPLGPRGMPGER
GRLGPQGAPGQRGAHGMPGKPGPMGPLGIPGSSGFPGNPGMKEAGPTGARGPEGPQGQRGETGPP
GPVGSPGLPGAIGTDGTPAKGPTGSPGTSGPPGSAGPPGSPGPQGSTGPQGIRGQPGDGVPGFK
GEAGPKGEPGPHGIQGPIGPPGEEGKRGPRGDPTVGPPGPVGERGAPGNRGFPGSDGLPGPKAQ
GERGPVGSSGPKGSGDPGRPGEPLPGARGLTGNPGVQGPEGKLGPLGAPGEDGRPGPPSIGIR
GQPGSMGLPGPKGSSGDPGPGEAGNAGVPGQRGAPGKDGEVGPSPVGPPGLAGERGEQGPPGPT
GFQGLPGPPGPPGEGGKPGDQGVPGDPAVGPLGPRGERGNPGERGEPGITGLPGEKGMAAGHGPD
GPKGSPGPSGTPGDTGPPGLQGMPGERGIAGTPGPKDRGGIGEKGAEGTAGNDGARGLPGPLGPP
GPAGPTGEKGEPGPRGLVGGPSRGNSRGENGPTGAVGFAGPQGPQGVKGEPGEPGQKGDA
GSPGPQGLAGSPGPQPHGPNGVPGLKGGRGTQGPPGATGFGPSAGRVPGPAGAPGPAGPLGEPGKE
GPPGLRGDPGSHGRVGDRGPAGPPGGPGDKGDPGEDQPGPDGPPGPAGTTGQRGIVGMPGQRGER
GMPGLPGPAGTPGKVGVPTGATGDKGPPGPVGPPGSNGPVGEPGPEGPAGNDGTPRDGAVGERGDR
GDPPGPAGLPGSQGAPGTPGPVGAPGDAGQRGDPGSRGPIGPPGRAGKRLGLPGQGPRGDKGDHGDR
GDRGQKGHRGFTGLQGLPGPPGPNGEQGSAGIPGPFGPRGPPGPVGPSGKEGNPGPLGPIGPPGVR
GSVGEAGPEGPPGEPGPPGPPGPPGHL

TAALGDIMGHYDESMPDPLPEFTEDQAAPDDKNKTDPGVHATLKSLSQIETMRSPDGSKKHPART
CDDLKLCHSAKQSGEYWIDPNQGSVEDAIKVYCNMETGETCISANPSSVPRKTWWASKSPDNKPVW
YGLDMNRGSQFAYGDHQSPNTAITQMTFLRLLSKEASQNITYICKNSVGYMDQAKNLKAVVLKG
ANDLDIKAEGNIRFRYIVLQDTCSKRNGNVGKTVFEYRTQNVARLPIIDLAPDVGGTDQEFGVVEI
GPVCFV

Col-V α3 sequence

MGNRRDLGQPRAGLCLLAALQLLPGTQADPVDVLKALGVQGGQAGVPEGPGFCPQRTPEGDRAFR
IGQASTLGIPTWELFPEGHFPENFSLITLRGQPANQSVLSSIYDERGARQLGLALGPALGLGDP
FRPLPQQVNLTDGRWHRVAVSIDGEMVTLVADCEAQPPVLGHGPRFISIAGLTVLGTQDLGEKTFE
GDIQELLISPDPQAAFQACERYLPDCDNLAPAATVAPQGEPETPRPRRKKGKGKRKKGRGRKGKGR
KKNKEIWTSPPPDSAENQTSTDIPKTETPAPNLPPPTPLVVTSTVTTGLNATILEGSLDPDSGT
ELGTLETKAAREDEEGDDSTMGPDFRAAEYPSRTQFOQIFPGAGEKGAKGEPAVIEKGQOFEGPPGA
PGPQGVVGPSGPPGPPGFPGDPGPPGPAGLPGIPGIDGIRGPPGTIVIMMPFQFAGGSFKGPPVSFQ
QAQAQAVLQQTQLSMK

GPPGPVGLTGRPGPVGLPGHPGLKGEEGAEGPQGPRGLQGPHGPPGRVGKMGRPGADGARGLPGDT

GPTGRPGVTGIDGAPGAKGNVGPPGEPGPPGQQGNHGSQGLPGPQGLIGTPGEKGPPGNPGIPGLP

GSDGPLGHPGHEGPTGEKGAQGPPGSAGPPGYPGPRGVKGTSGNRGLQGEKGEDGFPGFKGDV

GLKG DQG KPG APG PRGE DG PEG PKG QAG QAG E EG PPG SAGE KG KLG VP GL PG Y PGR PG PKG SIG FP

GPLGP¹IGEK²GKSGKTGQ³PGLERGPPG⁴SRG⁵ERGQ⁶PGATGQ⁷PGPKGDVG⁸QDGAPG⁹IPGEKGLPGLQ¹⁰

GPPGFPGPKGPPGHQGKDGRPGHPQRGELGFQGQTGPPGPAGVLGPQGKTGEVGPLGERGPPGPP

GPPGEQGLPGLEGREGAKGELGPPGPLGKEGPAGLRGFPPPKGGPGDPGPTGLKGDKGPPGPVGAN

GSPGERGPLGPAGGIGLPGQSGSEGPGVGPAGKKGSRGERGPPGPTKDGIPGPLGPLGPPGAAGPS

GEEGDKGDVGAPGHKGSKGDKGDAGPPGQPGIRGPAGHPGPPGADGAQGRGGPPGLFGQKGDDGVR

GFVGIVGPPGLQGLPGPPGEKGEVGDVGSMGPHGAPGPGRPQGPTGSEGTPGLPGGVGQPGAVGEK

GERGDAGDPGPGAPGI PGPKGDI GEKGDSGPAGPPGKKGPGEDGAKGSVGPTGLPGDLGPP

GDPGVSGIDGSPGEKGDPGDVGGPGPPGASGEPGAPGPGKRGPSGHMGREGREGEKGAKGEPGPD

GPPGRTGPMGARGPPGRVGPEGLRGI PGPVGEPGLLGAPGQMGP^PPGPLGPSGLPGLKGDTGPKGK

GHIGLIGLIGPPGEAGEKGKDQGLPGVQGPPGPKGDPGPPGPIGSLGHPGPPGVAGPLGQKGSKGSP

GSMGPRGDTGPAGPPGPGAP

AETIHTGBBBBBFVVPVPTPVVE

DCEIWIDINQCARDSTRTVCTTACCECTTDIRLEIVREASWKRIFCGWISITRKRRI SIV
DADGSPVN VVQLNFLKLLSATARQNFTYSCQNAAAWLDEATGDYSHSARFLGTNGEELSFNQTTAT
TVSVPQDGCR LRKGQT KTLFEFSSSRAGFLPLWDVAATDFGQTNQKFGFELGPVC FSS

Supplementary figure 1. Sequences of collagen α 1, α 2, and α 3. The Gly-Xaa-Yaa repeats are denoted in individual block. The red and blue triangles indicate the cleavage sites detected for the sample which is digested 60 min and 3min, respectively. The hydroxylproline positions in α 1(V) are shown in green.